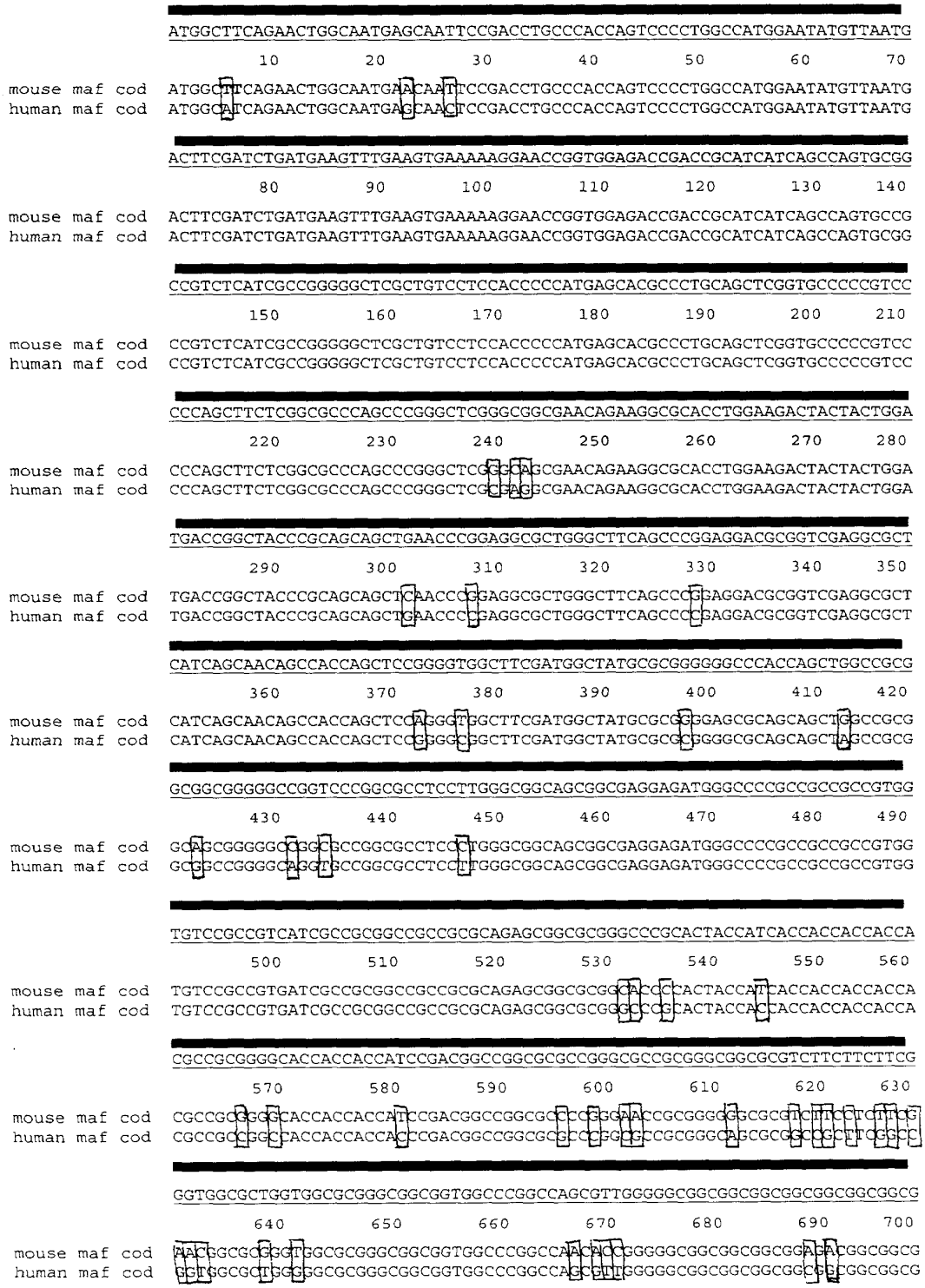
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mouse maf cod ACTTCGATCTGATGAAGTTTGAAGTGAAAAAGGAACCGGTGGAGACCGACCGCATCATCAGCCAGTGCCG
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mouse maf cod CCCAGCTTCTCGGCGCCCAGCCCGGGCTCGGGCGGCGAACAGAAGGCGCACCTGGAAGACTACTACTGGA
human maf cod CCCAGCTTCTCGGCGCCCAGCCCGGGCTCGGGCGGCGAACAGAAGGCGCACCTGGAAGACTACTACTGGA
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mouse maf cod TGACCGGCTACCCGCAGCAGCTCAACCCGGAGGCGCTGGGCTTCAGCCCGGAGGACGCGGTCGAGGCGCT
human maf cod TGACCGGCTACCCGCAGCAGCTGAACCCGGAGGCGCTGGGCTTCAGCCCGGAGGACGCGGTCGAGGCGCT
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mouse maf cod CATCAGCAACAGCCACCAGCTCCAGGGTGGCTTCGATGGCTATGCGCGGGGAGCGCAGCAGCTGGCCGCG
human maf cod CATCAGCAACAGCCACCAGCTCCGGGGCGGCTTCGATGGCTATGCGCGCGGGGCGCAGCAGCTAGCCGCG
GCGGCGGGGGCCGGTCCCGGCGCCTCCTTGGGCGGCAGCGGCGAGGAGATGGGCCCCGCCGCCGCCGTGG
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mouse maf cod GCAGCGGGGGCCGGCGCCGGCGCCTCCCTGGGCGGCAGCGGCGAGGAGATGGGCCCCGCCGCCGCCGTGG
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mouse maf cod CGCCGCGGGGCACCACCACCATCCGACGGCCGGCGCCCGGGAACCGCGGGGGGCGCGTCTTCCTCTTCG
human maf cod CGCCGCCGGCCACCACCACCACCCGACGGCCGGCGCGCCGGGCGCCGCGGGCAGCGCGGCCGCTTCGGCCC
GGTGGCGCTGGTGGCGCGGGCGGCGGTGGCCCGGCCAGCGTTGGGGGCGGCGGCGGCGGCGGCGGCGGCG
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mouse maf cod AACGGCGCGGGTGGCGCGGGCGGCGGTGGCCCGGCCAACACCGGGGGCGGCGGCGGCGGCGGCGGCG
human maf cod GGTGGCGCTGGTGGCGCGGGCGGCGGTGGCCCGGCCAGCGTTGGGGGCGGCGGCGGCGGCGGCGGCG

FIGURE 1A

GGGGCGGGGGGGGGCGGGGGGCGCCCTTCACCCGCACCATTCCGCGGGCGGCCTGCACTTCGACGACCG

710 720 730 740 750 760 770

mouse maf cod GGGGCACGGCGGCGGCGGGGGCGCCCTTCACCCGCACCATTCCGCGGGCGGCCTGCACTTCGACGACCG
human maf cod GAGGCGGCGGGGGGCGCGGCGGGCGCCCTGCACCCGCACCACGCCGCCGGCGGCCTGCACTTCGACGACCG

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780 790 800 810 820 830 840

mouse maf cod CTTCTCGGACGAGCAGTTGGTGACCATGTCGGTGCGCGAACTGAACCGGCAGCTGCGCGGGGTCAGCAAG
human maf cod CTTCTCCGACGAGCAGCTGGTGACCATGTCTGTGCGCGATTGAACCGGCAGCTGCGCGGGGTCAGCAAG

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850 860 870 880 890 900 910

mouse maf cod GAGGAGGTGATCCGACTGAAGCAGAAGAGGCGGACCCTGAAAAACCGCGGCTATGCCCAGTCCTGCCGCT
human maf cod GAGGAGGTGATCCGGCTGAAGCAGAAGAGGCGGACCCTGAAAAACCGCGGCTATGCCAAGTCCTGCCGCT

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920 930 940 950 960 970 980

mouse maf cod TCAAGAGGGTGCAGCAGAGACACGTCCTGGAGTCGGAGAAGAACCAGCTGCTGCAGCAGGTAGACCACCT
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990 1000 1010 1020 1030 1040 1050

mouse maf cod CAAGCAGGAGATCTCCAGGCTGGTGCGCGAAAGGGACGCCTACAAGGAGAAATACGAGAAGCTGGTGAGC
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1060 1070 1080 1090 1100 1110 1120

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human maf cod AGCGGCTTCCGAGAAAACGGCTCGAGCAGCGACAACCCGTCCTCTCCCGAGTTTTTCATAACTGAGCCCA

XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX

1130 1140 1150 1160 1170 1180 1190

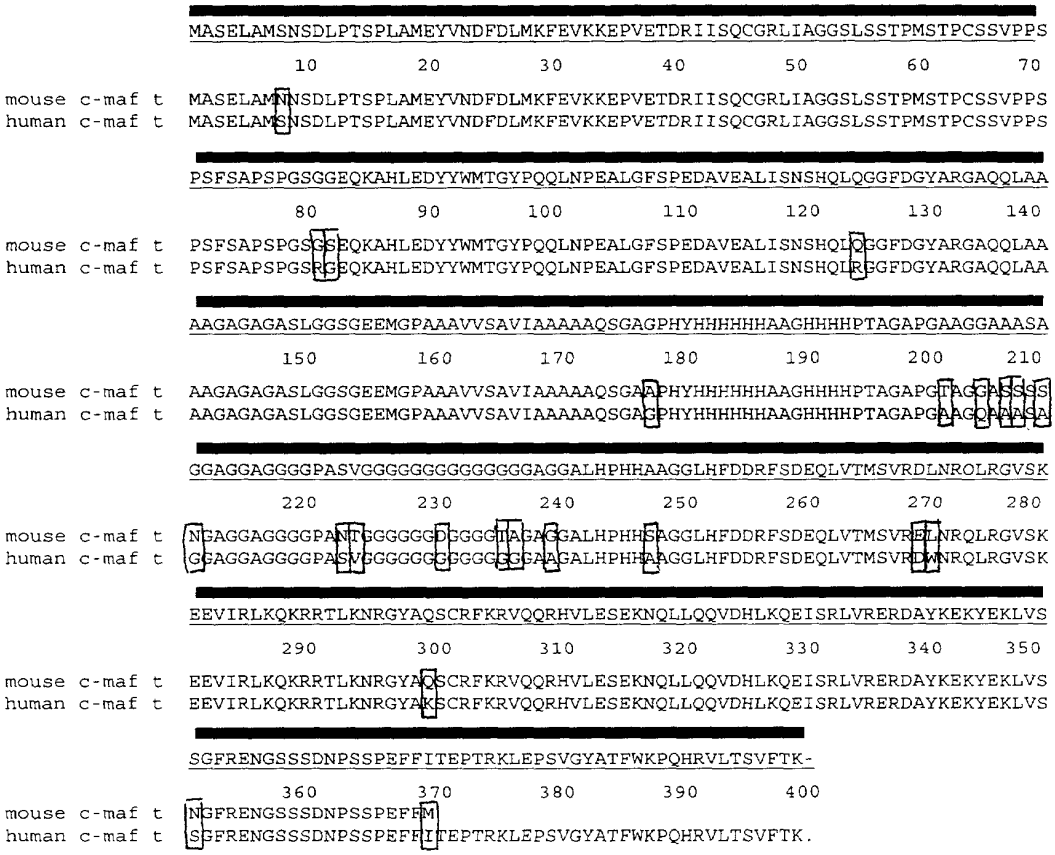
mouse maf cod
human maf cod CTCGCAAGTTGGAGCCATCAGTGGGATACGCCACATTTTGGAAGCCCCAGCATCGTGTACTTACCAGTGT

XXXXXXXXXXXXX

1200

mouse maf cod
human maf cod GTTCACAAAATGA

FIGURE 1B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mouse c-maf t MASELAMNNSDLPTSPLAMEYVNDFDLMKFEVKKEPVETDRIISQCGRLIAGGSLSSTPMSTPCSSVPPS
human c-maf t MASELAMSNSDLPTSPLAMEYVNDFDLMKFEVKKEPVETDRIISQCGRLIAGGSLSSTPMSTPCSSVPPS
PSFSAPSPGSGGEQKAHLEDYYWMTGYPQQLNPEALGFSPEDAVEALISNSHQLQGGFDGYARGAQQLAA
80 90 100 110 120 130 140
mouse c-maf t PSFSAPSPGSGSEQKAHLEDYYWMTGYPQQLNPEALGFSPEDAVEALISNSHQLQGGFDGYARGAQQLAA
human c-maf t PSFSAPSPGSRGEQKAHLEDYYWMTGYPQQLNPEALGFSPEDAVEALISNSHQLRGGFDGYARGAQQLAA
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mouse c-maf t AAGAGAGASLGGSGEEMGPAAAVVSAVIAAAAAQSGAAPHYHHHHHHAAGHHHHPTAGAPGTAGGASSSS
human c-maf t AAGAGAGASLGGSGEEMGPAAAVVSAVIAAAAAQSGAGPHYHHHHHHAAGHHHHPTAGAPGAAGGAAASA
GGAGGAGGGGPASVGGGGGGGGGGGGGAGGALHPHHAAGGLHFDDRFSDEQLVTMSVRDLNRQLRGVSK
220 230 240 250 260 270 280
mouse c-maf t NGAGGAGGGGPANTGGGGGGDGGGGTAGAGGALHPHHSAGGLHFDDRFSDEQLVTMSVRELNRQLRGVSK
human c-maf t GGAGGAGGGGPASVGGGGGGGGGGGGGAAGALHPHHAAGGLHFDDRFSDEQLVTMSVRDMNRQLRGVSK
EEVIRLKQKRRTLKNRGYAQSCRFKRVQQRHVLESEKNQLLQQVDHLKQEISRLVRERDAYKEKYEKLVS
290 300 310 320 330 340 350
mouse c-maf t EEVIRLKQKRRTLKNRGYAQSCRFKRVQQRHVLESEKNQLLQQVDHLKQEISRLVRERDAYKEKYEKLVS
human c-maf t EEVIRLKQKRRTLKNRGYAKSCRFKRVQQRHVLESEKNQLLQQVDHLKQEISRLVRERDAYKEKYEKLVS
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360 370 380 390 400
mouse c-maf t NGFRENGSSSDNPSSPEFFM
human c-maf t SGFRENGSSSDNPSSPEFFITEPTRKLEPSVGYATFWKPQHRVLTSVFTK.

FIGURE 2